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RAW SEQUENCE LISTING   DATE: 10/12/2001
PATENT APPLICATION: US/09/896,720   TIME: 11:17:38

Input Set : A:\5853173u.app
Output Set: N:\CRF3\10122001\I896720.raw

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    3 <110> APPLICANT: SONG, WEN-YUAN
    4        PI, LI-YA
    6 <120> TITLE OF INVENTION: UBIQUITIN LIGASE
    8 <130> FILE REFERENCE: 5853-173
   10 <140> CURRENT APPLICATION NUMBER: 09/896,720
C-->11 <141> CURRENT FILING DATE: 2001-09-24
   13 <150> PRIOR APPLICATION NUMBER: 60/215,049
   14 <151> PRIOR FILING DATE: 2000-06-29
   16 <160> NUMBER OF SEQ ID NOS: 24
   18 <170> SOFTWARE: PatentIn Ver. 2.1
   20 <210> SEQ ID NO: 1
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   22 <212> TYPE: DNA
   23 <213> ORGANISM: Oryza sativa
   25 <400> SEQUENCE: 1
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   27 cagctcggcg acctcgacgc cctggccgcc ctcctcgccg ccgacccttc cctcgctcgc 120
   28 cgcgccaccc tctacgaccg cctctccgtc ctccacatcg ccgccgccaa tggccgcatc 180
   29 gaggtgctct ccatgttctt ggatcgcggg gcgccgccgg acgcggtgaa tcggcacaag 240
   30 cagacgccgc tgatgctcgc ggccatgcac ggcaagatcg actgcgtgct caagctcctc 300
   31 caggccgacg caaatatctt gatgttcgac tcggtgcacg cgaggacctg cctccaccac 360
   32 gcggcgtact acggccacgt cgactgcctg caggccatcc tcgccgccgc gcagaccacg 420
   33 ccggtggccg actcatgggg tttcgcccgg ttcgtcaacg tcagggacga ccacggcgcc 480
   34 actccgctgc atctcgcggc caggcagggg cggccggggt gcgtgcaggt gttgctggag 540
   35 aacggcgcca ttgtgtcggc tttgacagga tcatatggct tccctggaag cacgtcgctt 600
   36 catttggctg ctcgtagcgg gaacttggat tgcatcagga agctgcttgc ctggggagct 660
   37 gatcggctcc aaagggattc ggctgggaga attccctatt ctgttgcgct gaaacggaac 720
   38 catggagcat gtgcagcttt gctgaaccct acatcagcag agcccatggt gtggccatcc 780
   39 ccacttaagt tcatcagtga gcttgaacca gaagctaagg ctctcctgga agcagctctg 840
   40 atggaagcca cagggagag ggagaagaaa atcctgaatg cacaaagta ctccctgcca 900
   41 tccccttcgc ccggtgatga cagtgccgat gacgatgcat gctcagaggt gagcgacacg 960
   42 gagctttgct gcatctgctt cgaccaggct tgcaccattg aggtgcaaga ctgtggacat 1020
   43 caaatgtgtg caccgtgcac gctggcactg tgctgtcaca acaaacccaa tccgacgacc 1080
   44 ctgacaccgc cctcaccggc ctgcccattc tgccggggca gcatctcacg gctggtggtg 1140
   45 gcccaaacaa ggtctgcttg tgatcctgac aagccgtcat ccctgcagct cacccggaag 1200
   46 cggtcgcgtc gatctcacaa cctcagtgag ggcagcagca gcttcaaagg gctaccttcg 1260
   47 gccatgggct ccttctcaaa gcttggccgt ggctcgagcc gcatggcgga cagtgacagc 1320
   48 agcaacctgg acaagcctga gcacgatcta tga                          1353
   51 <210> SEQ ID NO: 2
   52 <211> LENGTH: 450
   53 <212> TYPE: PRT
   54 <213> ORGANISM: Oryza sativa
   56 <220> FEATURE:
   57 <221> NAME/KEY: MOD_RES
   58 <222> LOCATION: (258)
   59 <223> OTHER INFORMATION: Any amino acid
   61 <400> SEQUENCE: 2
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62 Met Gly His Gly Val Ser Cys Ala Arg Thr Gly Asp Glu His Asp Phe
63   1               5               10              15
65 Phe Arg Ala Ala His Leu Gly Asp Leu Asp Ala Leu Ala Ala Leu Leu
66           20              25              30
68 Ala Ala Asp Pro Ser Leu Ala Arg Arg Ala Thr Leu Tyr Asp Arg Leu
69           35              40              45
71 Ser Val Leu His Ile Ala Ala Ala Asn Gly Arg Ile Glu Val Leu Ser
72       50              55              60
74 Met Phe Leu Asp Arg Gly Ala Pro Pro Asp Ala Val Asn Arg His Lys
75  65              70              75              80
77 Gln Thr Pro Leu Met Leu Ala Ala Met His Gly Lys Ile Asp Cys Val
78               85              90              95
80 Leu Lys Leu Leu Gln Ala Asp Ala Asn Ile Leu Met Phe Asp Ser Val
81           100             105             110
83 His Ala Arg Thr Cys Leu His His Ala Ala Tyr Tyr Gly His Val Asp
84           115             120             125
86 Cys Leu Gln Ala Ile Leu Ala Ala Ala Gln Thr Thr Pro Val Ala Asp
87       130             135             140
89 Ser Trp Gly Phe Ala Arg Phe Val Asn Val Arg Asp Asp His Gly Ala
90 145             150             155             160
92 Thr Pro Leu His Leu Ala Ala Arg Gln Gly Arg Pro Gly Cys Val Gln
93           165             170             175
95 Val Leu Leu Glu Asn Gly Ala Ile Val Ser Ala Leu Thr Gly Ser Tyr
96           180             185             190
98 Gly Phe Pro Gly Ser Thr Ser Leu His Leu Ala Ala Arg Ser Gly Asn
99       195             200             205
101 Leu Asp Cys Ile Arg Lys Leu Leu Ala Trp Gly Ala Asp Arg Leu Gln
102     210             215             220
104 Arg Asp Ser Ala Gly Arg Ile Pro Tyr Ser Val Ala Leu Lys Arg Asn
105 225             230             235             240
107 His Gly Ala Cys Ala Ala Leu Leu Asn Pro Thr Ser Ala Glu Pro Met
108             245             250             255
110 Val Xaa Pro Ser Pro Leu Lys Phe Ile Ser Glu Leu Glu Pro Glu Ala
111         260             265             270
113 Lys Ala Leu Leu Glu Ala Ala Leu Met Glu Ala Asn Arg Glu Arg Glu
114         275             280             285
116 Lys Lys Ile Leu Asn Gly Thr Lys Tyr Ser Leu Pro Ser Pro Ser Pro
117     290             295             300
119 Gly Asp Asp Ser Ala Asp Asp Asp Ala Cys Ser Glu Val Ser Asp Thr
120 305             310             315             320
122 Glu Leu Cys Cys Ile Cys Phe Asp Gln Ala Cys Thr Ile Glu Val Gln
123             325             330             335
125 Asp Cys Gly His Gln Met Cys Ala Pro Cys Thr Leu Ala Leu Cys Cys
126             340             345             350
128 His Asn Lys Pro Asn Pro Thr Thr Leu Thr Pro Pro Ser Pro Ala Cys
129         355             360             365
131 Pro Phe Cys Arg Gly Ser Ile Ser Arg Leu Val Val Ala Gln Thr Arg
132         370             375             380
134 Ser Ala Cys Asp Pro Asp Lys Pro Ser Ser Leu Gln Leu Thr Arg Lys
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W--> (marking at line 110)

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```
135 385                   390                   395                   400
137 Arg Ser Arg Arg Ser His Asn Leu Ser Glu Gly Ser Ser Ser Phe Lys
138               405                   410                   415
140 Gly Leu Pro Ser Ala Met Gly Ser Phe Ser Lys Leu Gly Arg Gly Ser
141               420                   425                   430
143 Ser Arg Met Ala Asp Ser Asp Ser Ser Asn Leu Asp Lys Pro Glu His
144           435                   440                   445
146 Asp Leu
147       450
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152 <212> TYPE: DNA
153 <213> ORGANISM: Artificial Sequence
155 <220> FEATURE:
156 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
158 <400> SEQUENCE: 3
159 gtcgaccaga tctcataaga gaagaaaga                            29
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163 <211> LENGTH: 31
164 <212> TYPE: DNA
165 <213> ORGANISM: Artificial Sequence
167 <220> FEATURE:
168 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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171 gtcgacgaat gaaatcagtt gtgaagggta c                        31
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175 <211> LENGTH: 35
176 <212> TYPE: DNA
177 <213> ORGANISM: Artificial Sequence
179 <220> FEATURE:
180 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
182 <400> SEQUENCE: 5
183 ggatccgtcg accacaagag aactaaaaag ggagc                    35
186 <210> SEQ ID NO: 6
187 <211> LENGTH: 42
188 <212> TYPE: DNA
189 <213> ORGANISM: Artificial Sequence
191 <220> FEATURE:
192 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
194 <400> SEQUENCE: 6
195 ggatccgtcg accccgggca gaagtcgatc tgaagtgtgg ca            42
198 <210> SEQ ID NO: 7
199 <211> LENGTH: 30
200 <212> TYPE: DNA
201 <213> ORGANISM: Artificial Sequence
203 <220> FEATURE:
204 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
206 <400> SEQUENCE: 7
207 ggatccatga tatccatggg tcacggtgtc                          30
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Input Set : A:\5853173u.app
Output Set: N:\CRF3\10122001\I896720.raw

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211 <211> LENGTH: 31
212 <212> TYPE: DNA
213 <213> ORGANISM: Artificial Sequence
215 <220> FEATURE:
216 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
218 <400> SEQUENCE: 8
219 cgggatccga tatcagatgc agcaaagctc c                             31
222 <210> SEQ ID NO: 9
223 <211> LENGTH: 30
224 <212> TYPE: DNA
225 <213> ORGANISM: Artificial Sequence
227 <220> FEATURE:
228 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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235 <211> LENGTH: 24
236 <212> TYPE: DNA
237 <213> ORGANISM: Artificial Sequence
239 <220> FEATURE:
240 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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247 <211> LENGTH: 30
248 <212> TYPE: DNA
249 <213> ORGANISM: Artificial Sequence
251 <220> FEATURE:
252 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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255 ggatccatga tatccgatgc atgctcagag                                30
258 <210> SEQ ID NO: 12
259 <211> LENGTH: 30
260 <212> TYPE: DNA
261 <213> ORGANISM: Artificial Sequence
263 <220> FEATURE:
264 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
266 <400> SEQUENCE: 12
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271 <211> LENGTH: 28
272 <212> TYPE: DNA
273 <213> ORGANISM: Artificial Sequence
275 <220> FEATURE:
276 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
278 <400> SEQUENCE: 13
279 gaattctcta gaccggggca gcatctca                                  28
282 <210> SEQ ID NO: 14
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284 <212> TYPE: DNA
285 <213> ORGANISM: Artificial Sequence
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290 <400> SEQUENCE: 14
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294 <210> SEQ ID NO: 15
295 <211> LENGTH: 28
296 <212> TYPE: DNA
297 <213> ORGANISM: Artificial Sequence
299 <220> FEATURE:
300 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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303 gaattcagat ctccggggca gcatctca                               28
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312 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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318 <210> SEQ ID NO: 17
319 <211> LENGTH: 17
320 <212> TYPE: DNA
321 <213> ORGANISM: Artificial Sequence
323 <220> FEATURE:
324 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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331 <211> LENGTH: 17
332 <212> TYPE: DNA
333 <213> ORGANISM: Artificial Sequence
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336 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
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339 ggtgcatgct ccaatgg                                           17
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343 <211> LENGTH: 21
344 <212> TYPE: DNA
345 <213> ORGANISM: Artificial Sequence
347 <220> FEATURE:
348 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
350 <400> SEQUENCE: 19
351 gcgctgaaac ggaaccatgg a                                      21
354 <210> SEQ ID NO: 20
355 <211> LENGTH: 22
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VERIFICATION SUMMARY   DATE: 10/12/2001
PATENT APPLICATION: US/09/896,720   TIME: 11:17:39

Input Set : A:\5853173u.app
Output Set: N:\CRF3\10122001\I896720.raw

L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:110 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2